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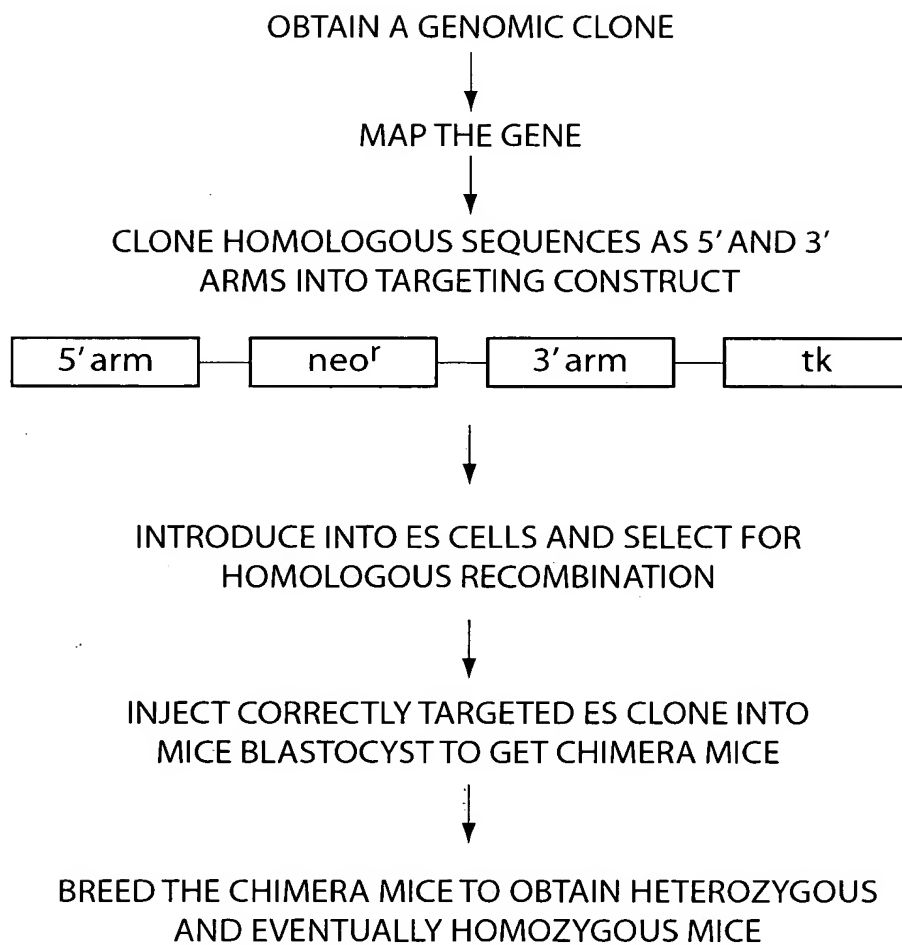


Fig. 1A

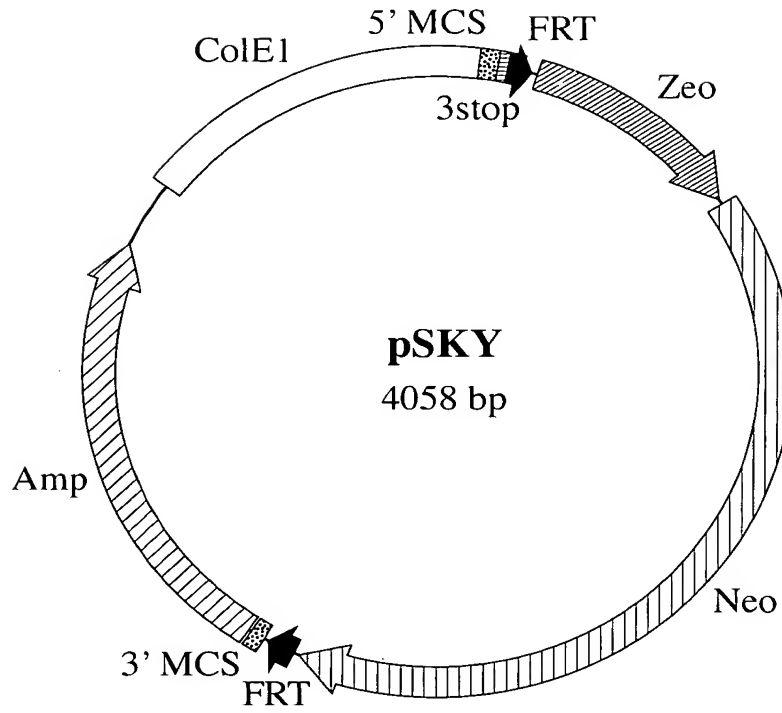


Fig. 1B

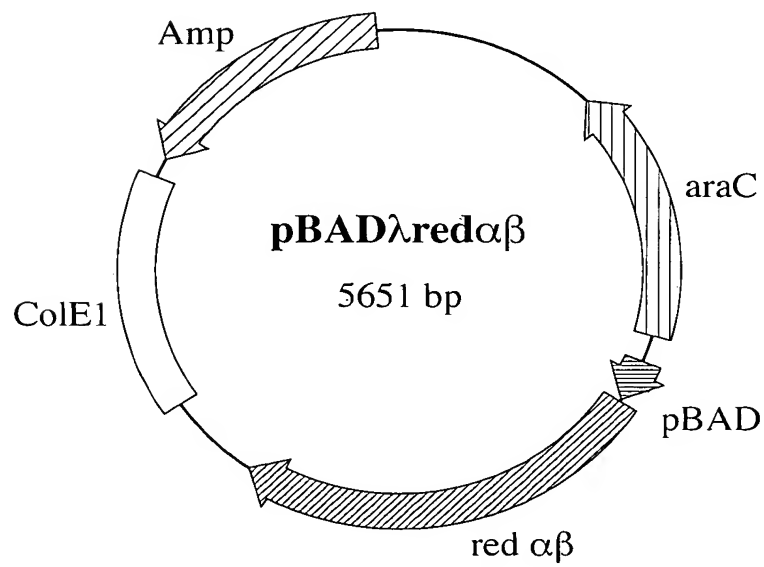


Fig. 1C

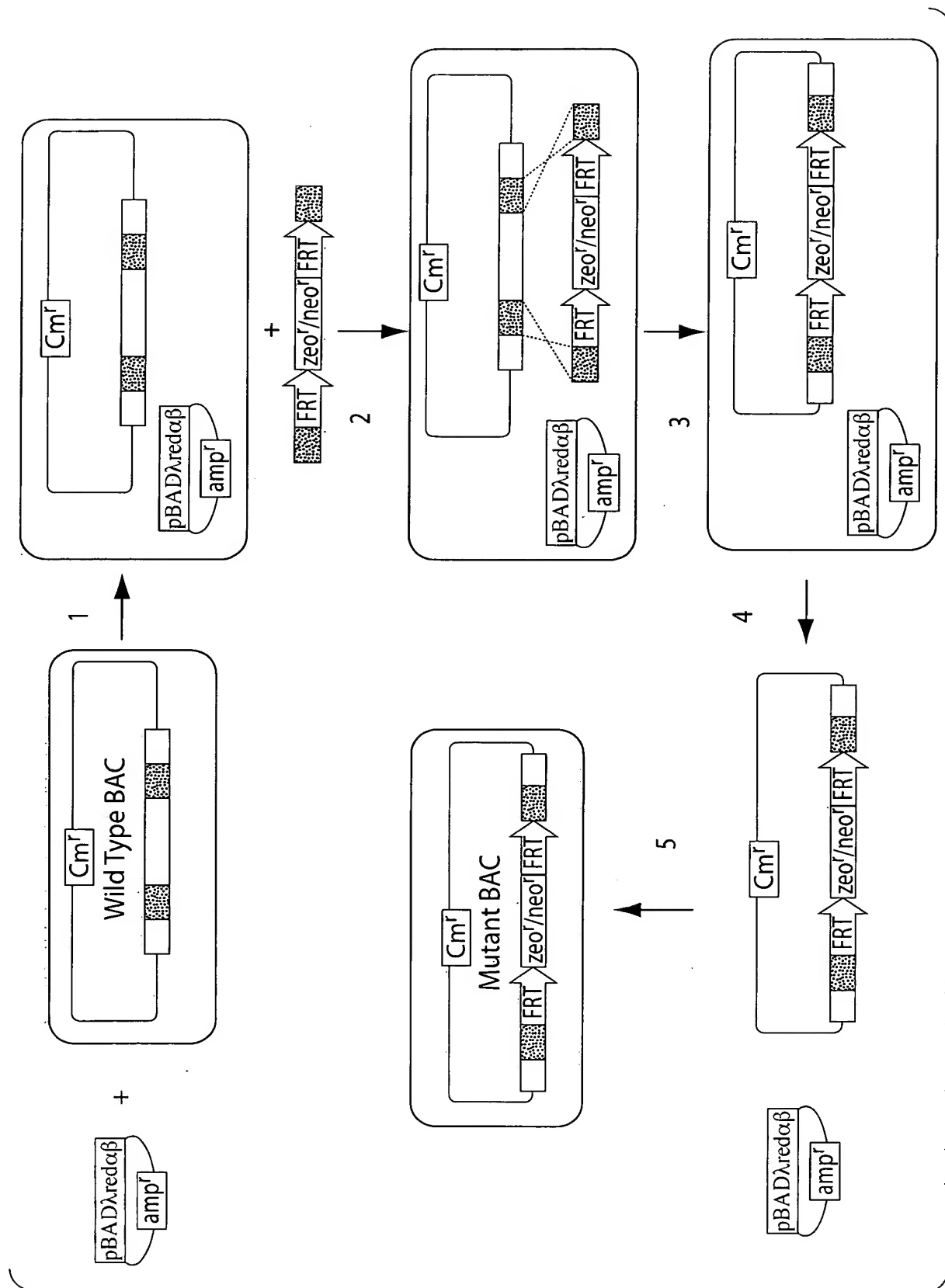


Fig. 1D

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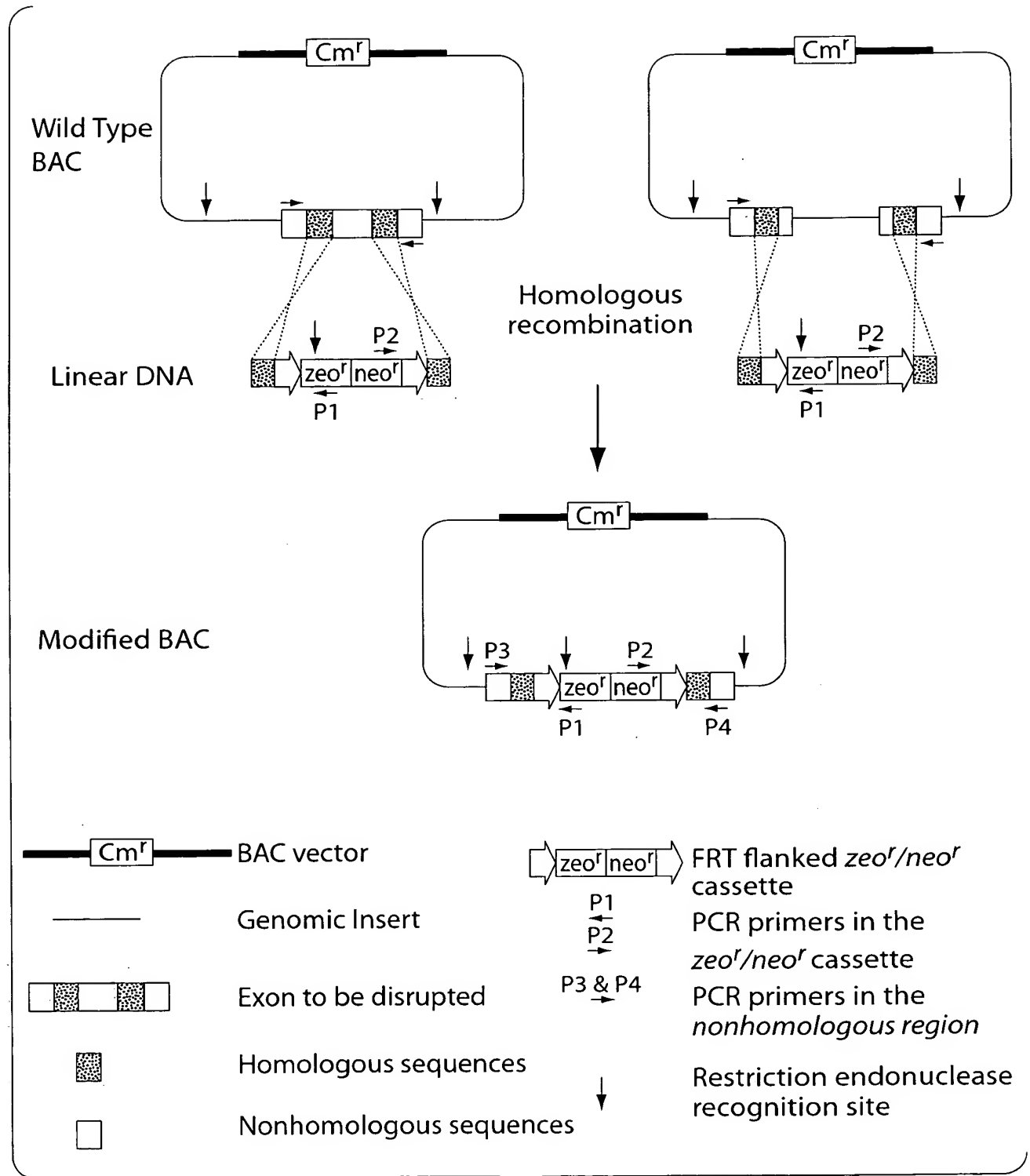


Fig. 1E

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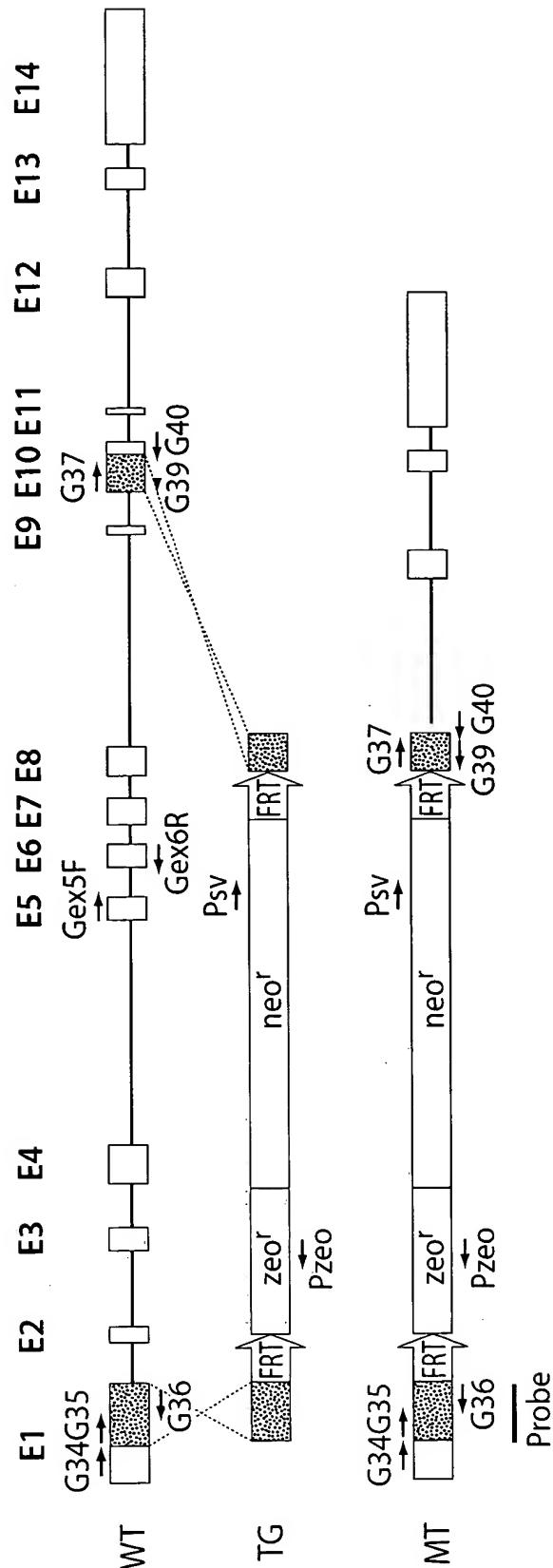


Fig. 2A

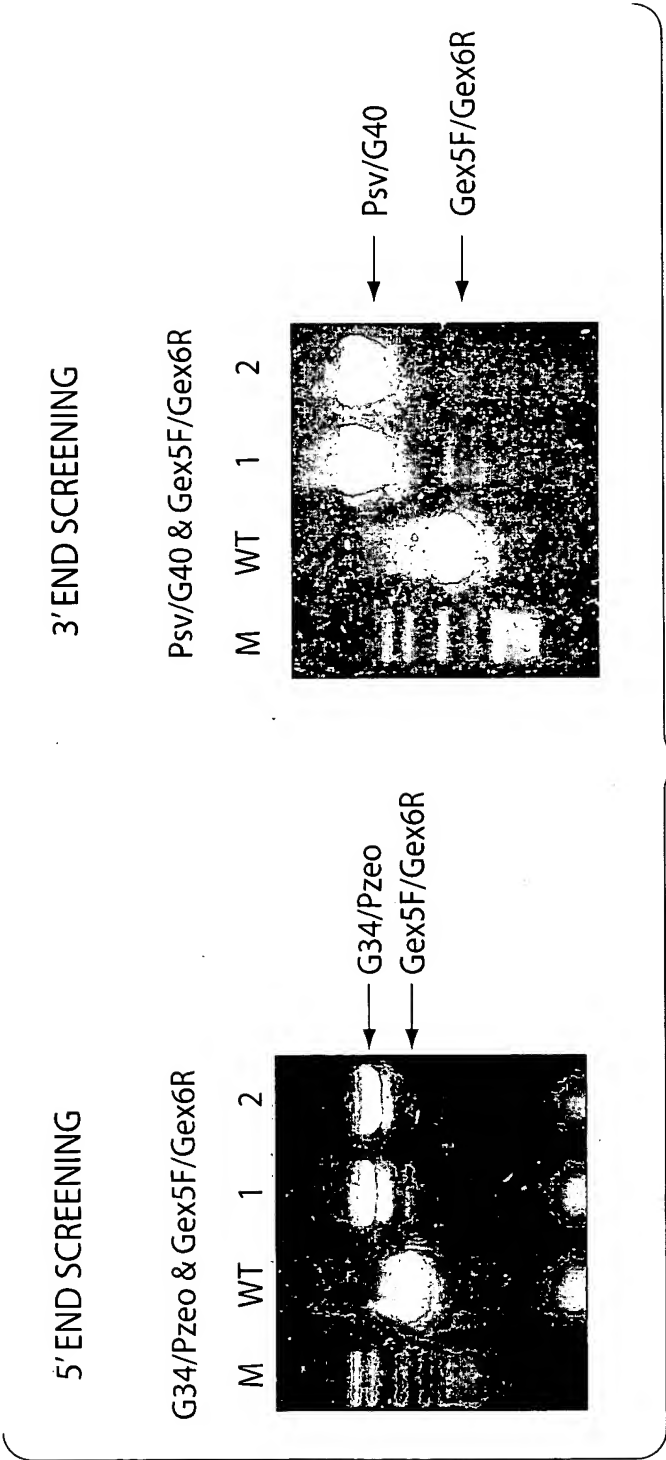


Fig. 2B

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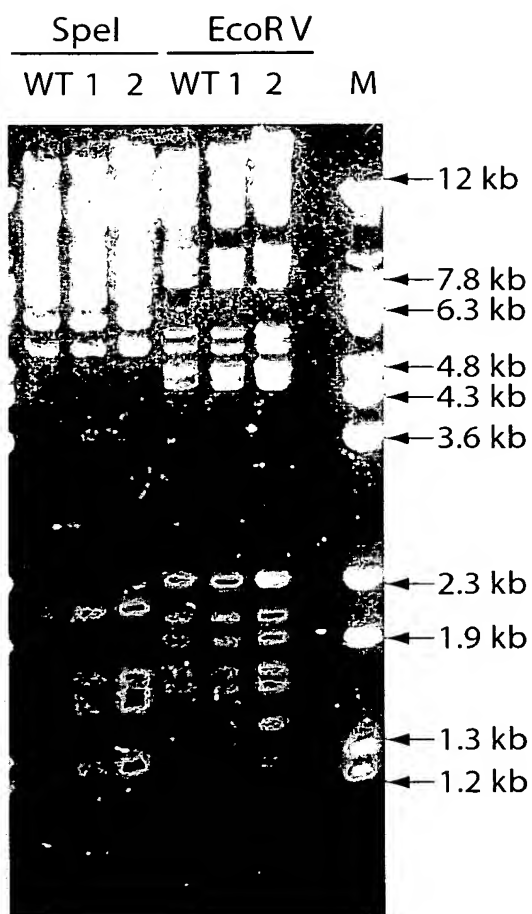


Fig. 2C

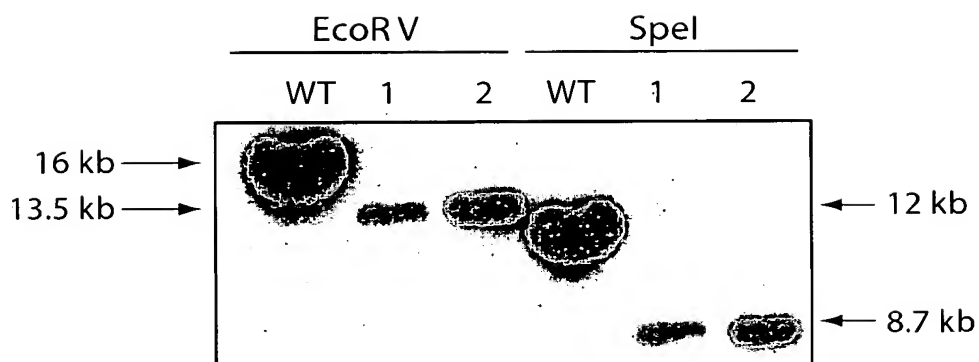


Fig. 2D

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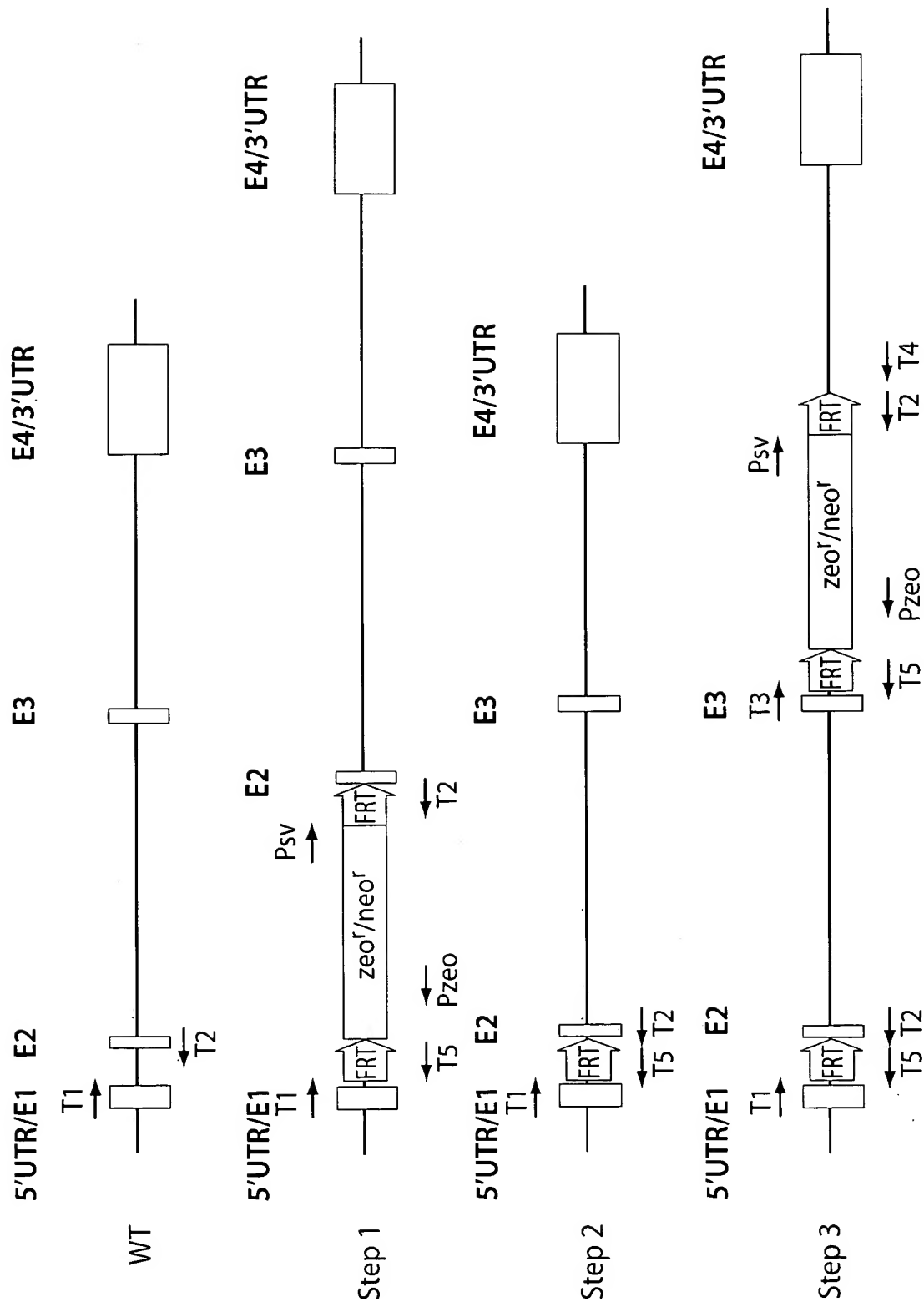


Fig. 3A

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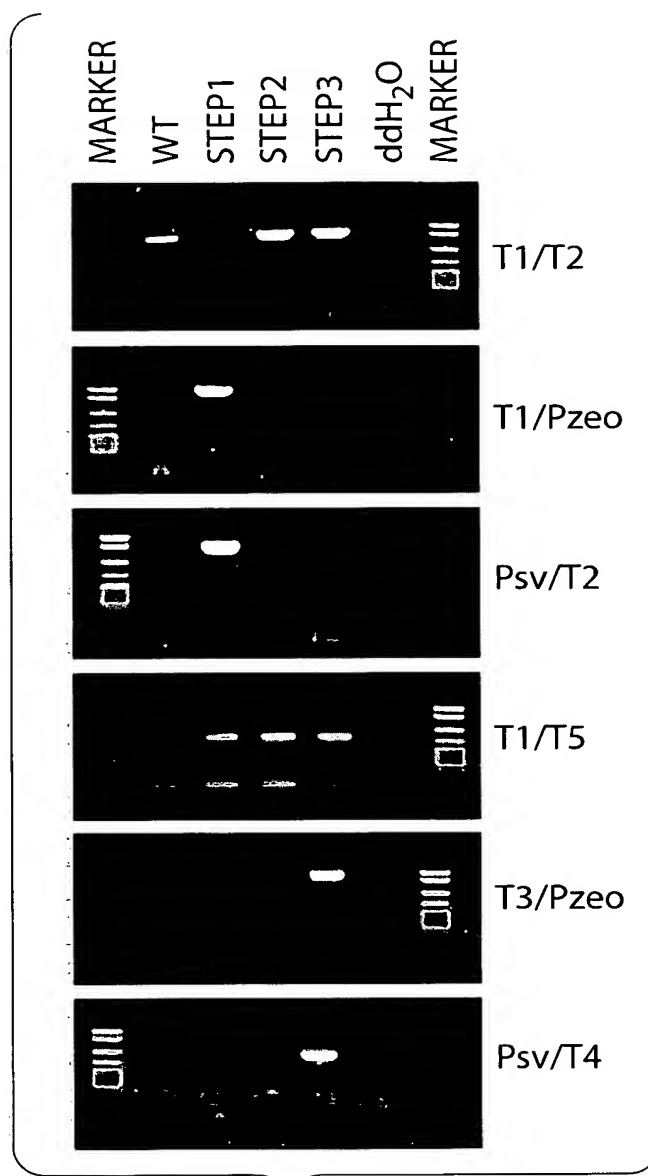
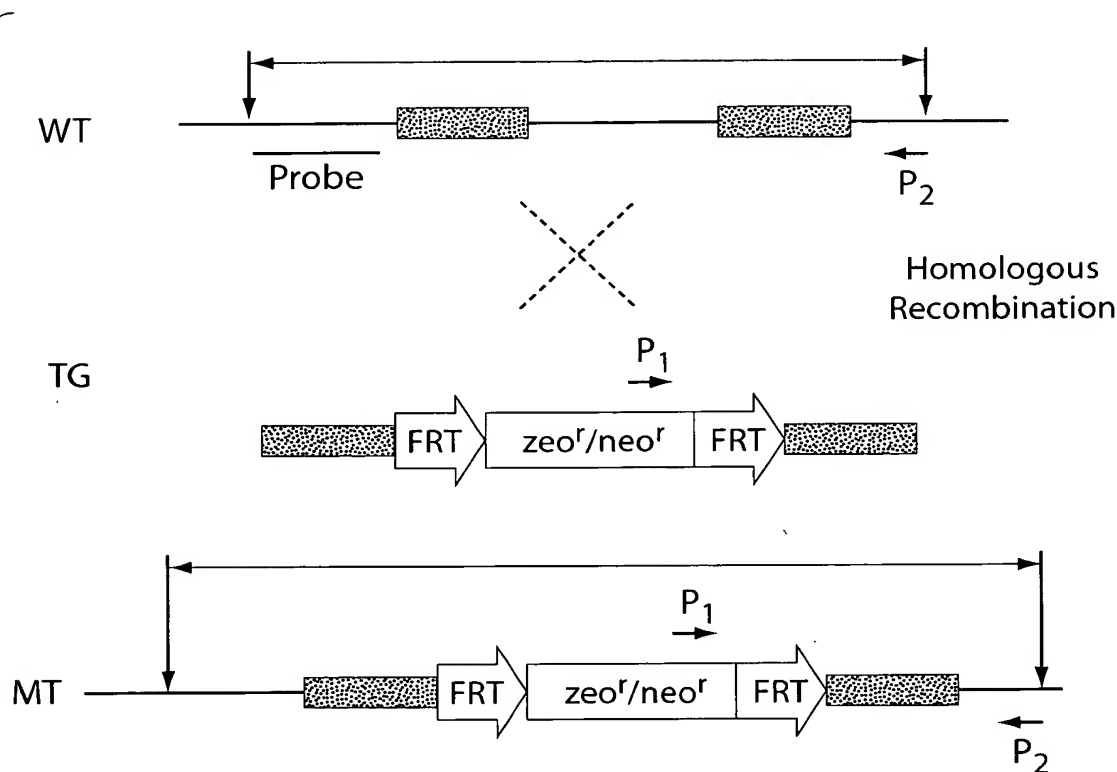


Fig. 3B

REPLACEMENT SHEET

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ASSAY	RESULTS
Genomic PCR w/ P1/P2	————
Southern Blotting	———— MT ———— WT
Genomic PCR w/ P1/P2	No Product
Southern Blotting	———— WT

Random
Insertion

Fig. 4A

REPLACEMENT SHEET

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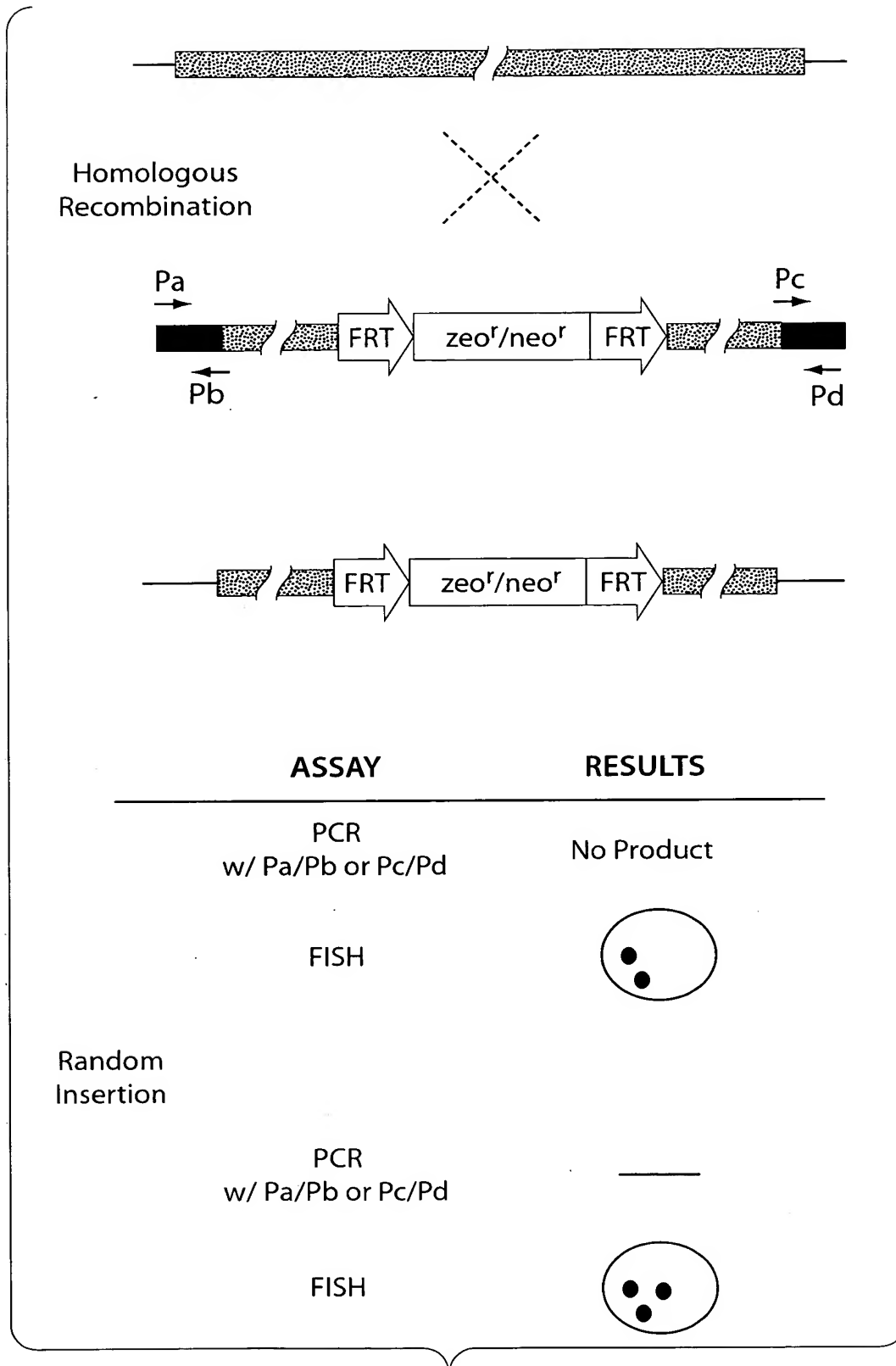


Fig. 4B

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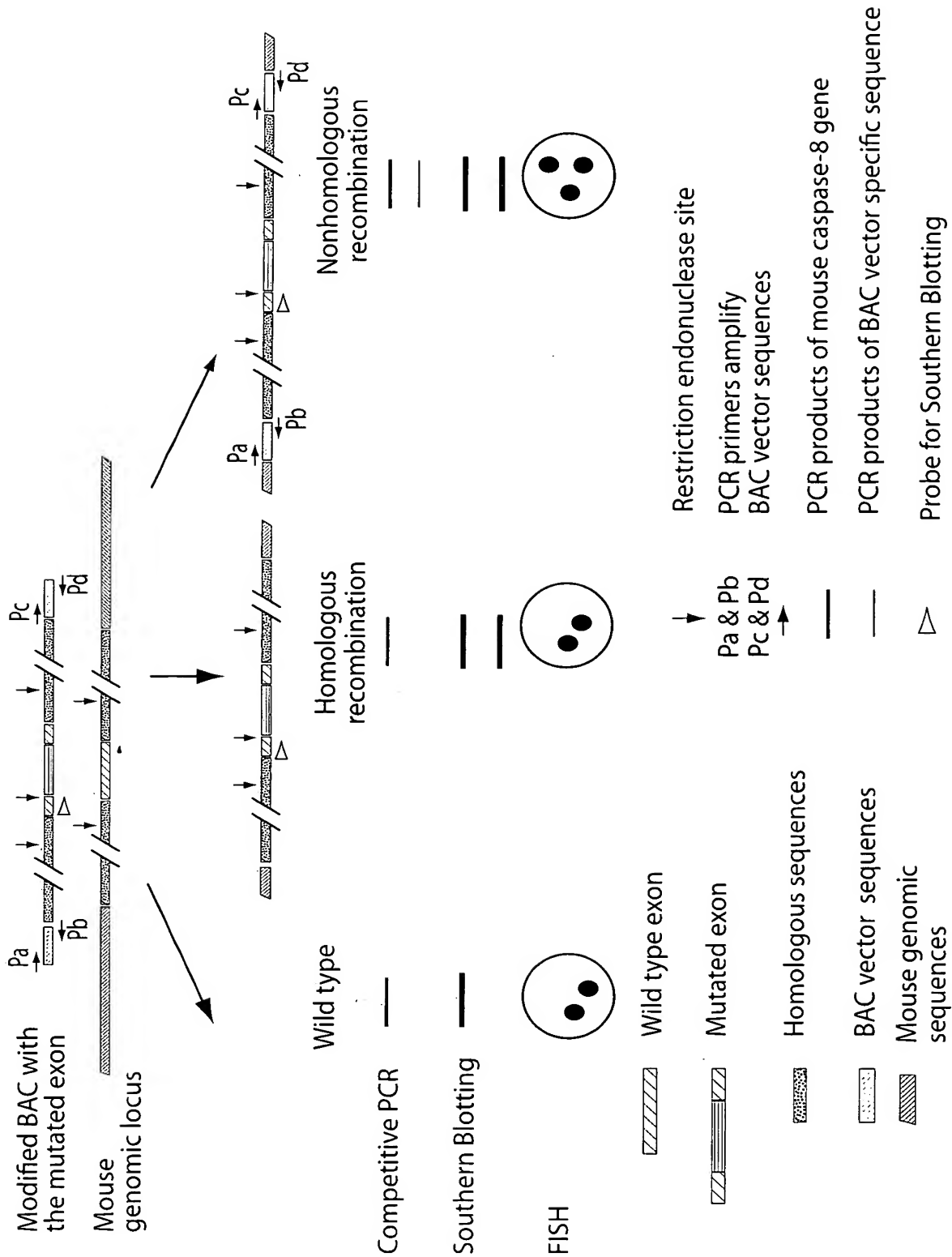


Fig. 4C

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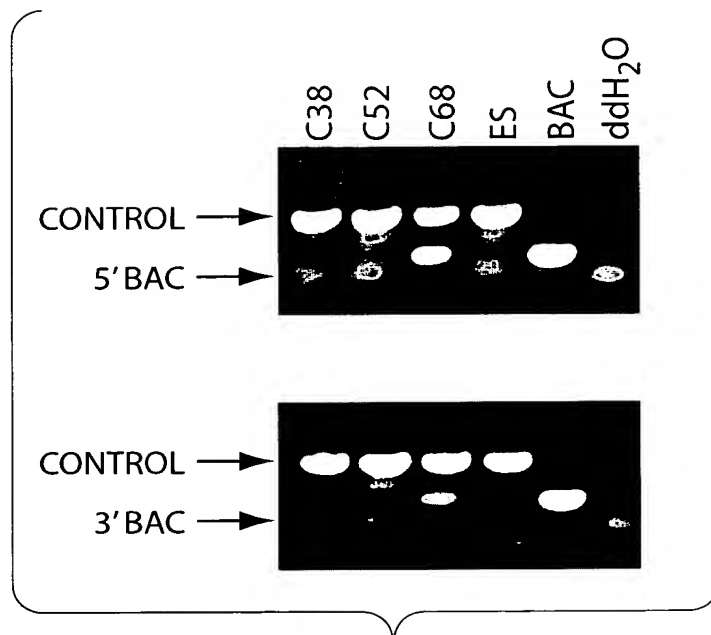


Fig. 5A

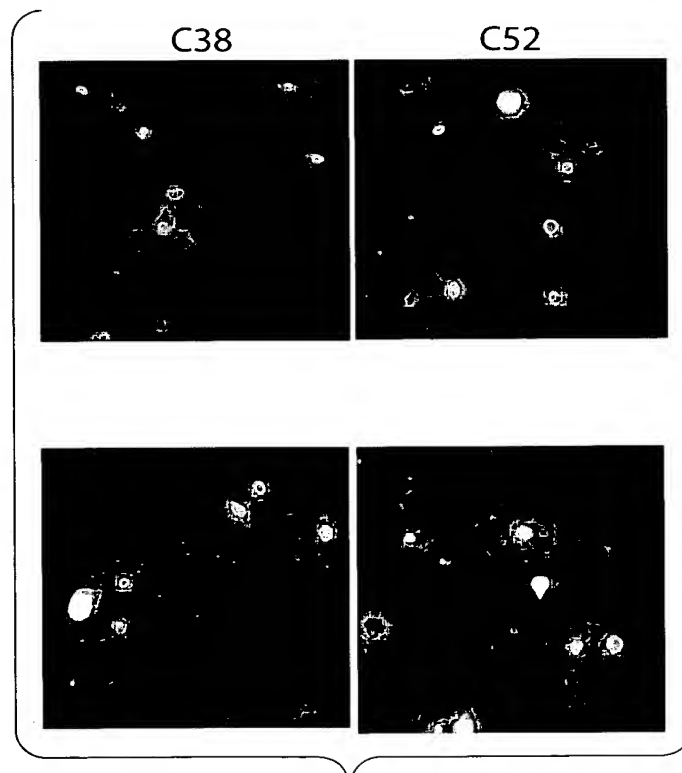


Fig. 5B

REPLACEMENT SHEET

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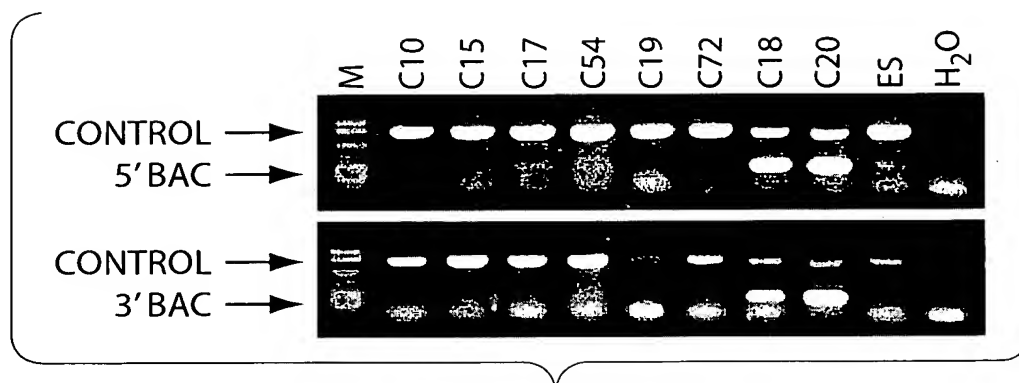


Fig. 6A

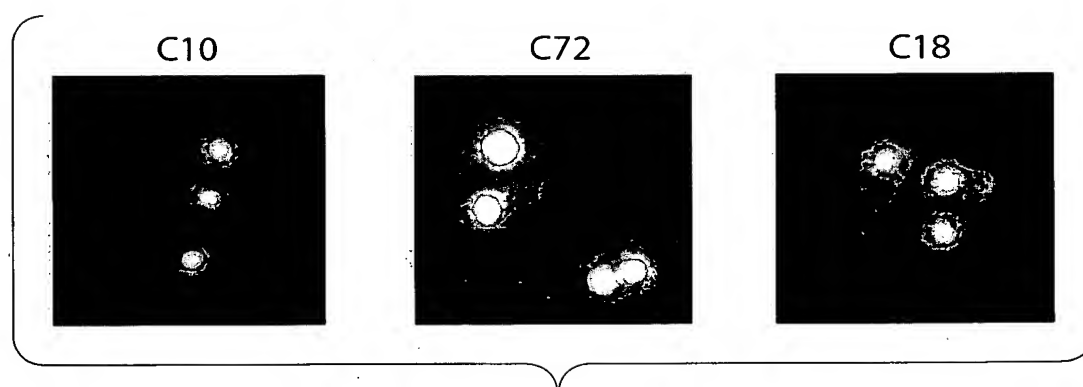


Fig. 6B

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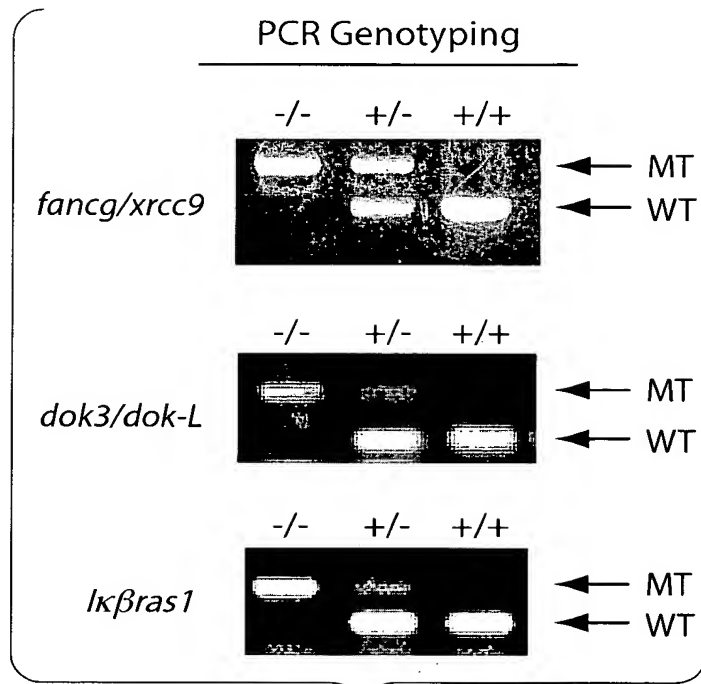


Fig. 7A

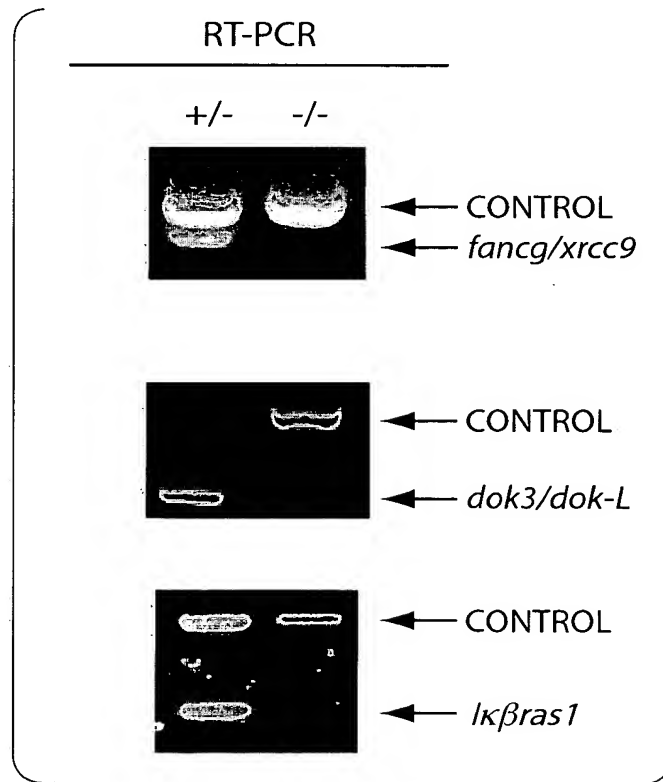


Fig. 7B

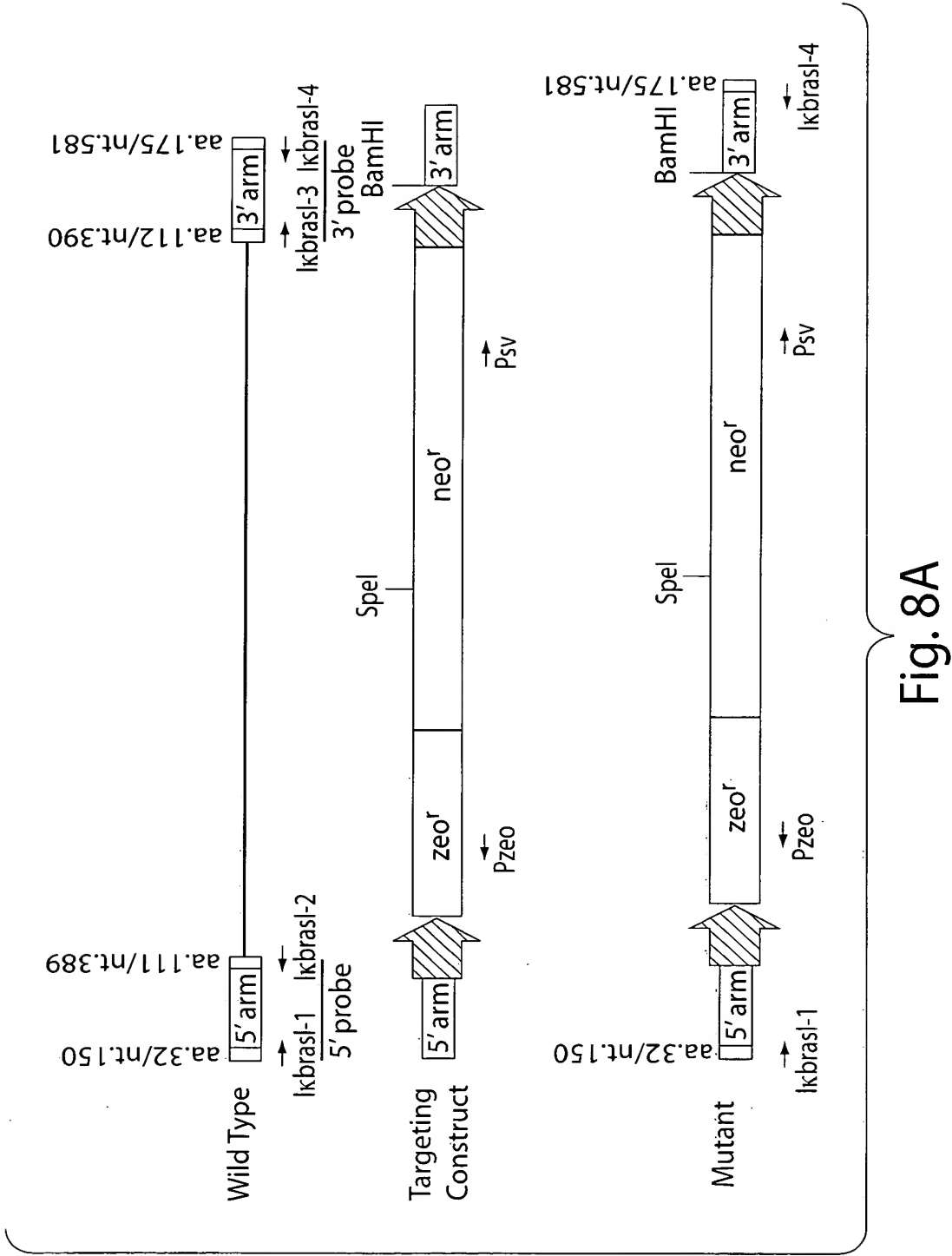


Fig. 8A

REPLACEMENT SHEET

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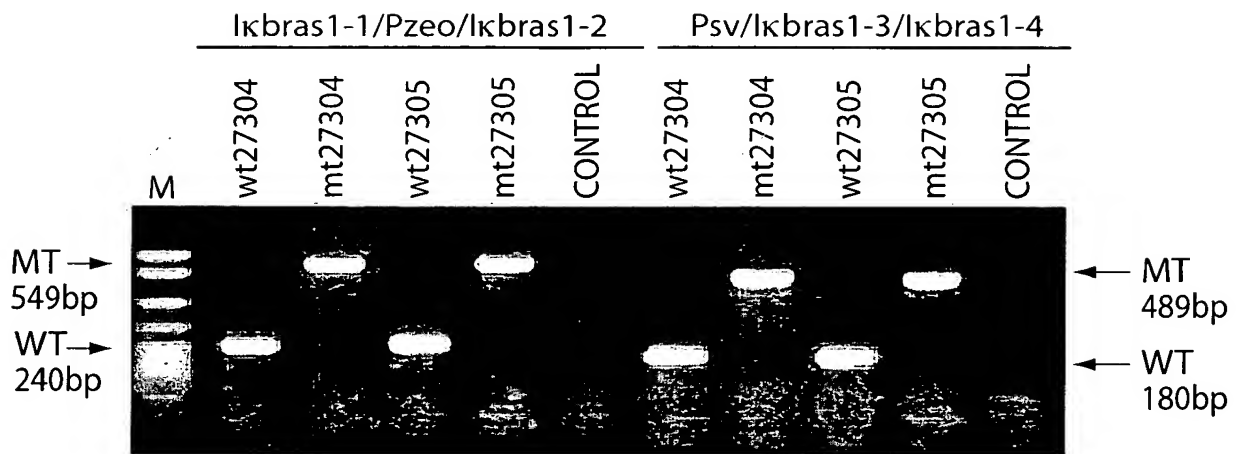


Fig. 8B

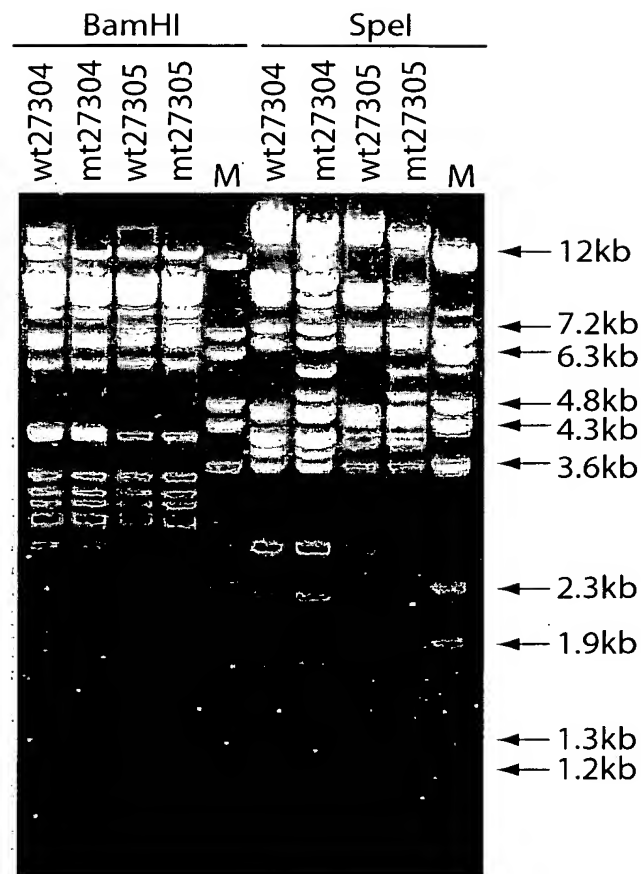


Fig. 8C

REPLACEMENT SHEET

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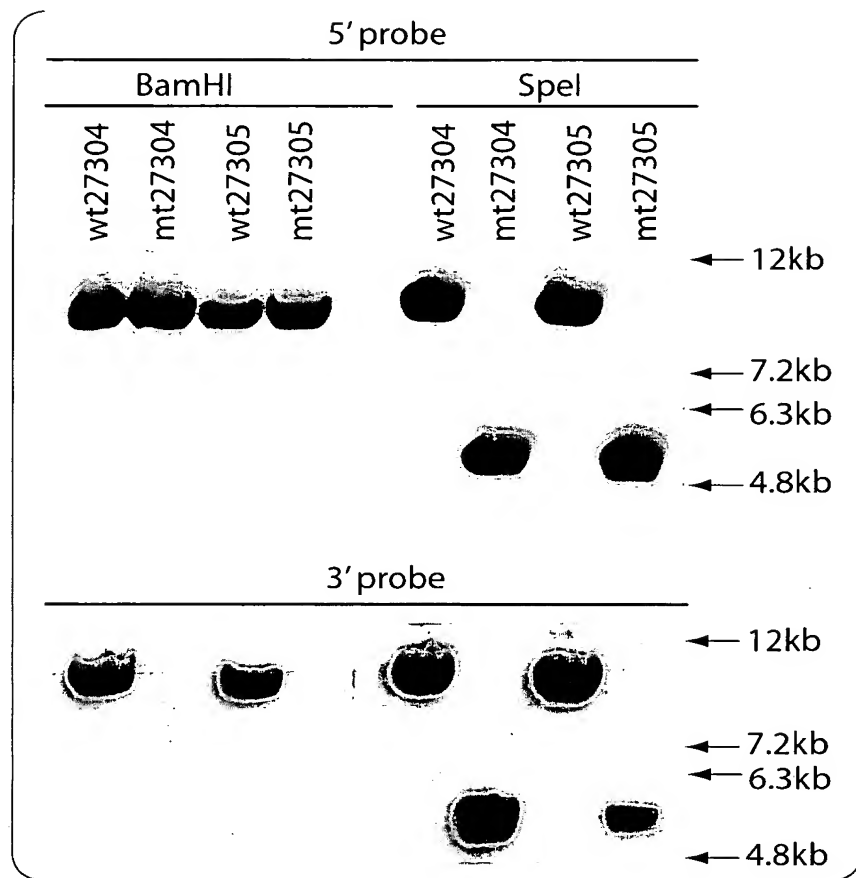


Fig. 8D

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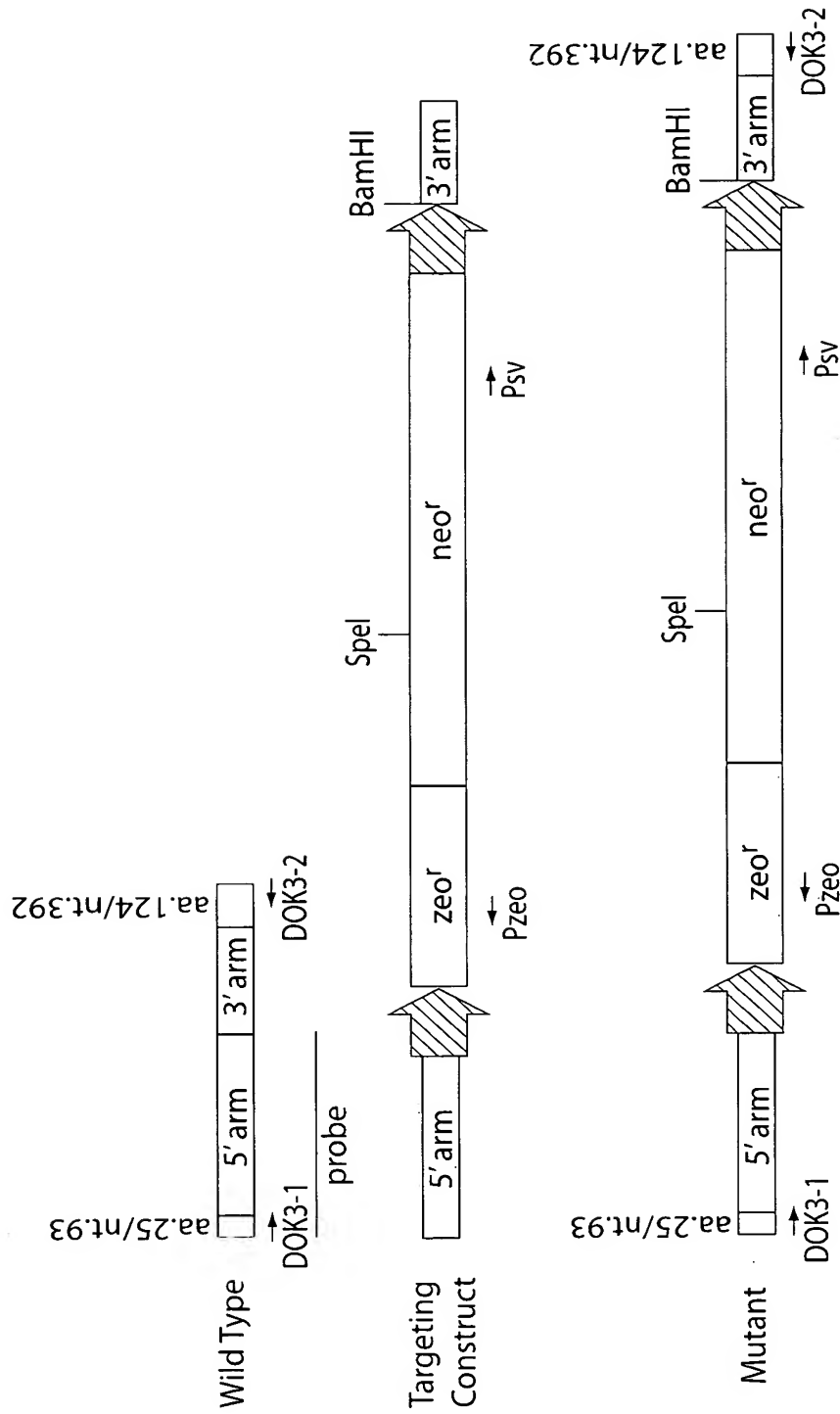


Fig. 9A

REPLACEMENT SHEET

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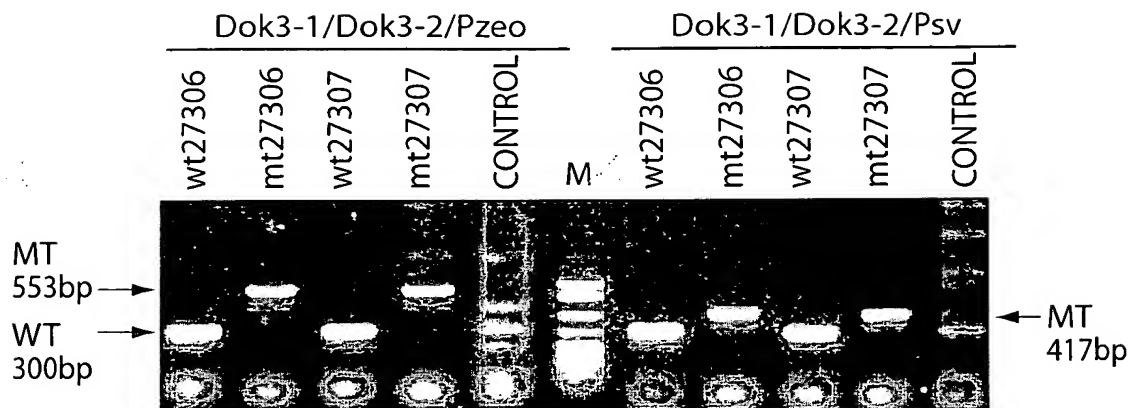


Fig. 9B

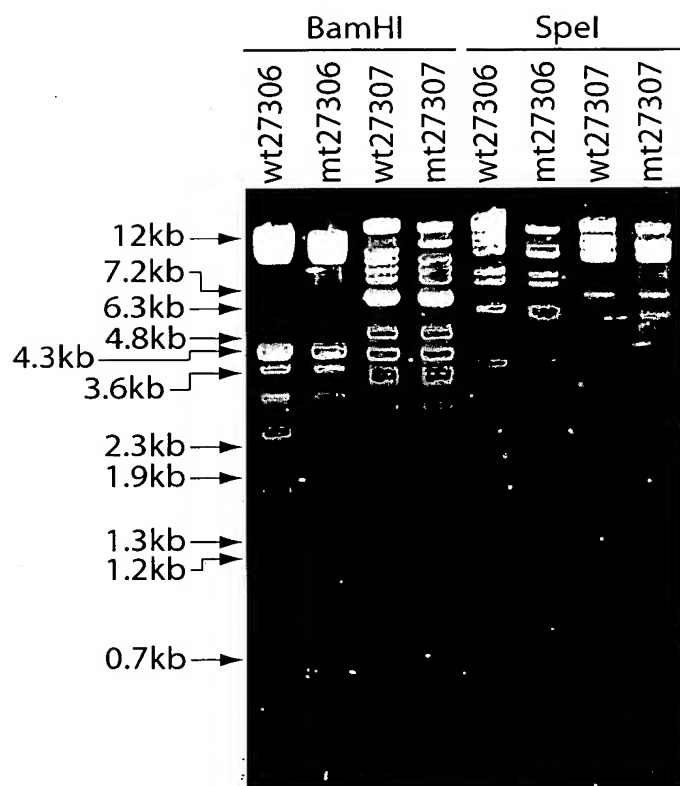


Fig. 9C

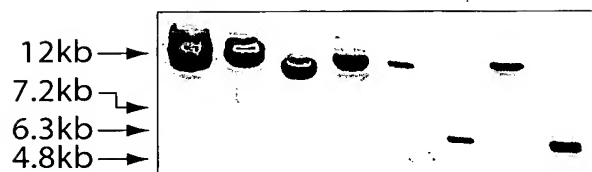


Fig. 9D

REPLACEMENT SHEET

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Gene/Clone	5'arm (nt)	3'arm (nt)	Deletion (nt)	Efficiency*
<i>fancg/xrcc9</i>	480	260	4900	100%
<i>Iκβras1.#1</i>	185	138	unknown	30%
<i>Iκβras1.#2</i>	185	138	unknown	25%
<i>tab2.#1</i>	213	340	869	30%
<i>tab2.#2</i>	213	340	869	50%
<i>pag.#1</i>	105	373	no	100%
<i>pag.#2</i>	105	373	no	58%
<i>pag.#3</i>	105	373	no	100%
<i>dok3/dok-L#1</i>	188	93	no	42%
<i>dok3/dok-L#2</i>	188	93	no	30%
<i>bruce</i>	50	50	no	100%

*The efficiency was calculated based on nested PCR screening of 24 ampicillin and chloramphenicol double resistant colonies at the 3' integration site.

Fig. 10

REPLACEMENT SHEET

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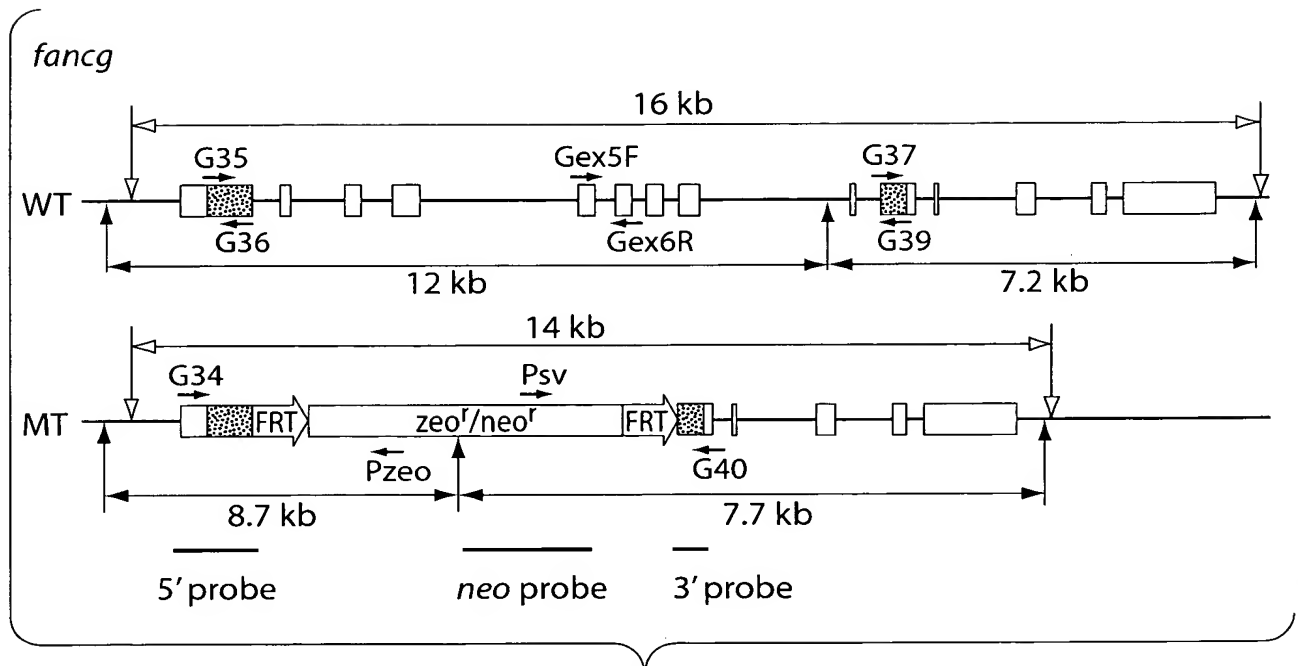


Fig. 11A

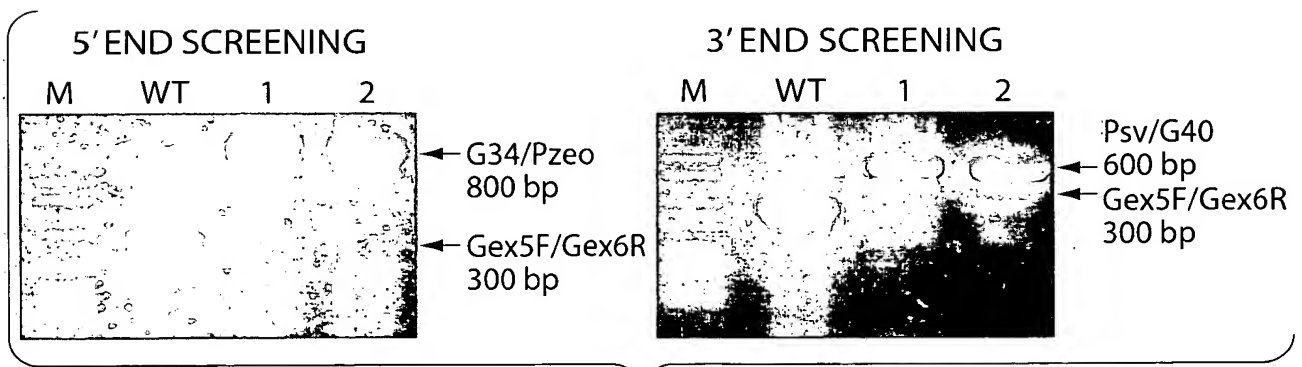


Fig. 11B

REPLACEMENT SHEET

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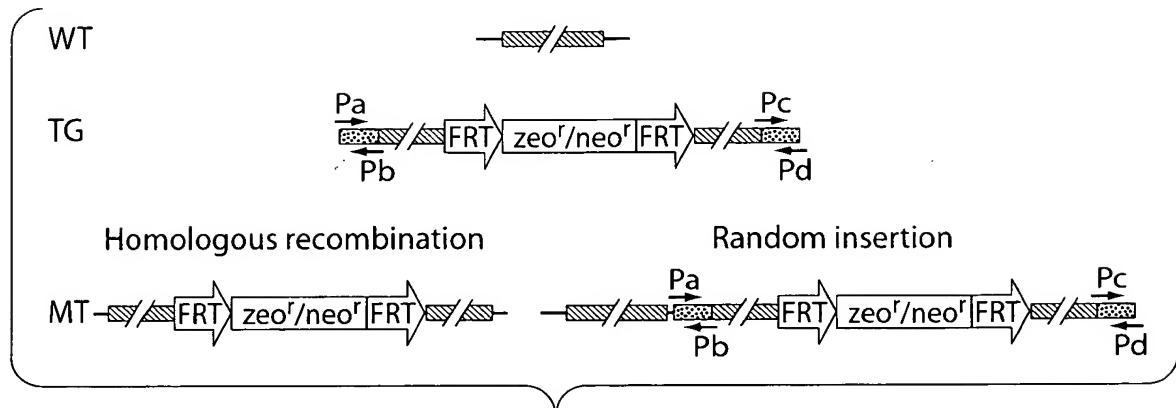


Fig. 12A

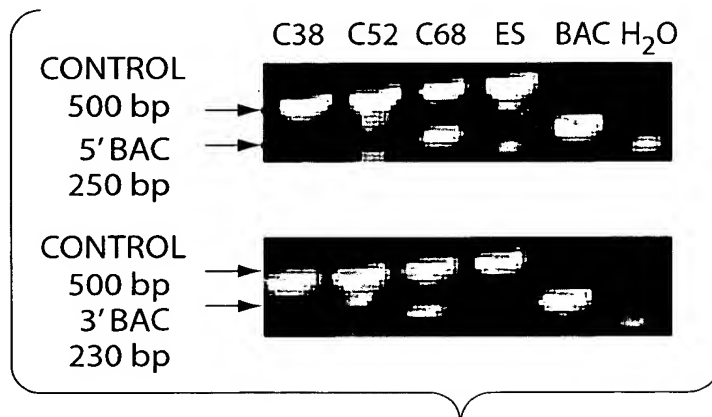


Fig. 12B

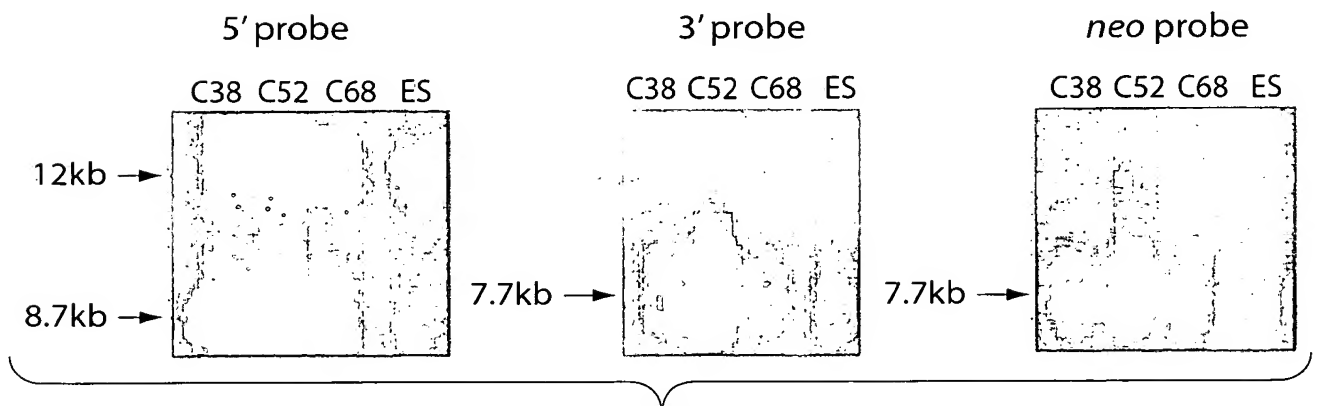


Fig. 12C

REPLACEMENT SHEET

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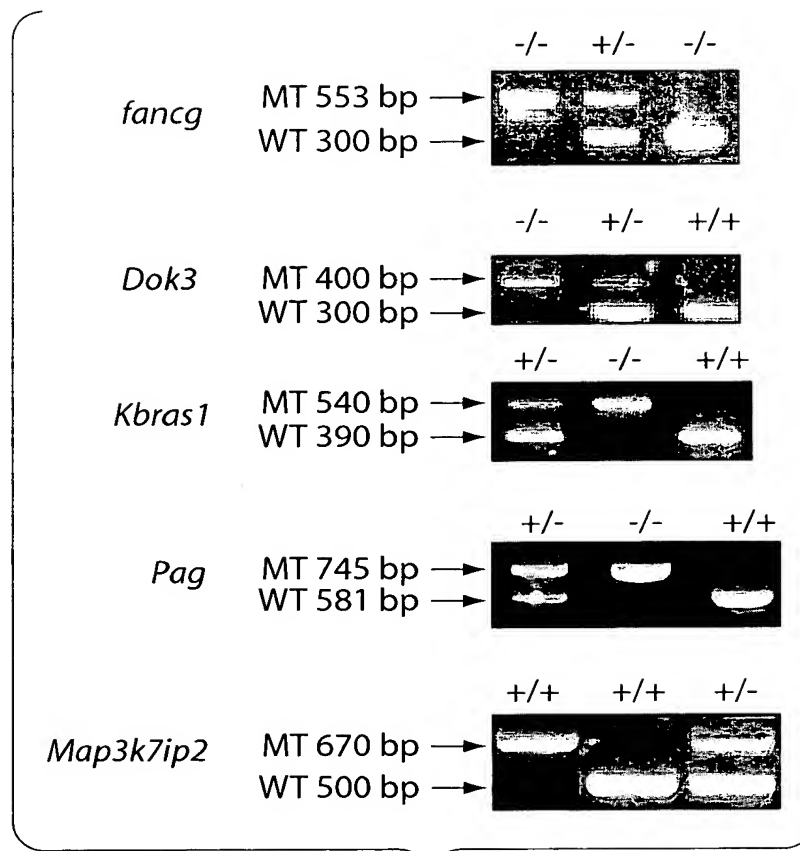


Fig. 13A

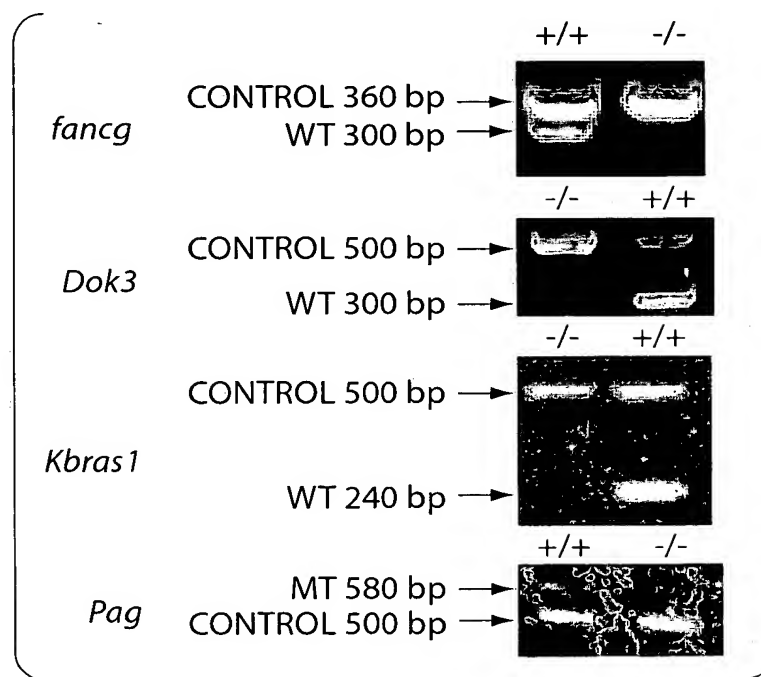


Fig. 13B

REPLACEMENT SHEET

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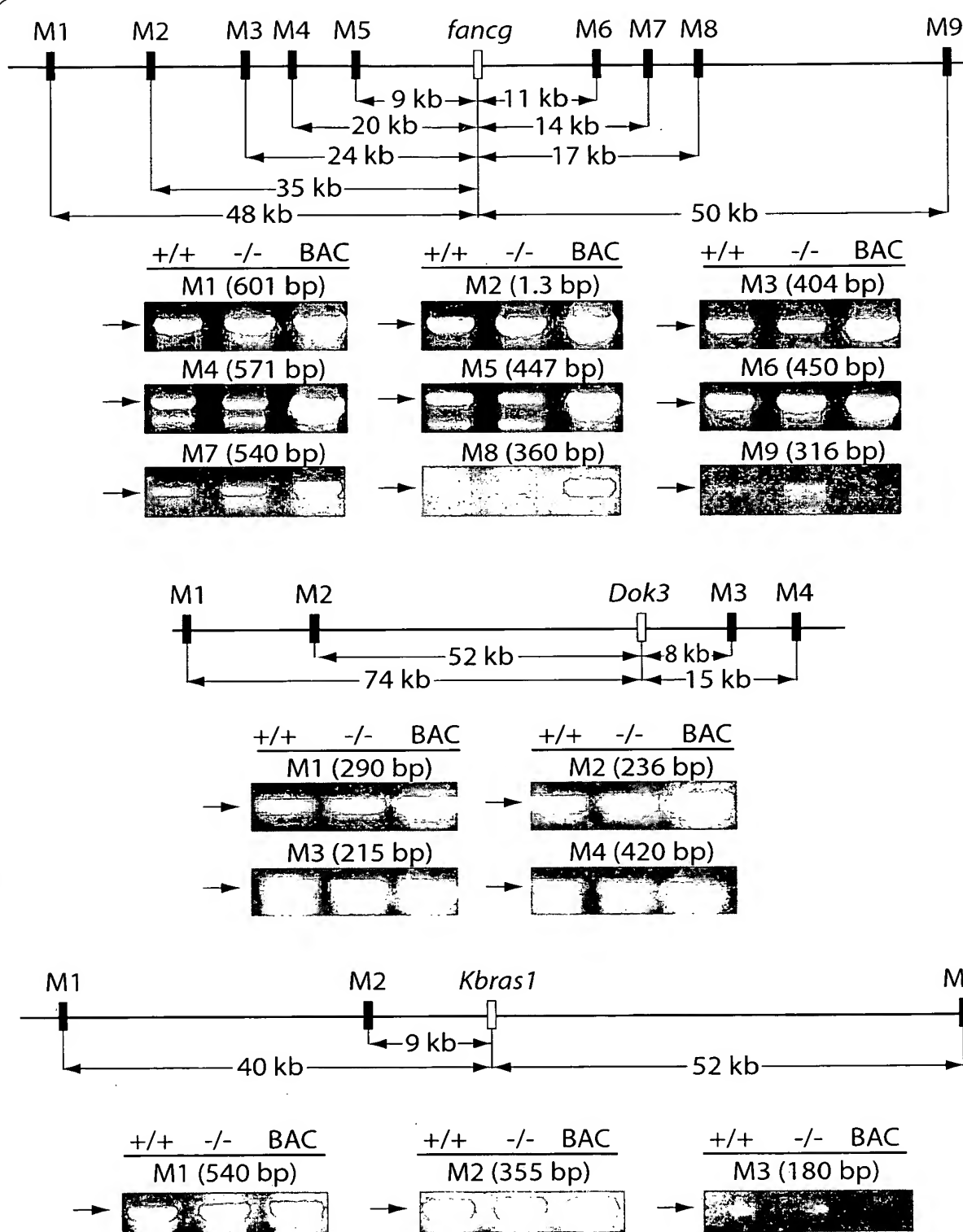


Fig. 13C

Gene	PCR screening/96 colonies					FISH screening			Effective targeting efficiency ^c
	no signal ^a	5' BAC	3' BAC	5' & 3' BAC	no BAC signal ^b	colonies screened	2 spots	3 spots	
<i>FANCG/XRCC9</i>	36	4	6	28	22 (37%)	4	3 (75%)	1	28%
<i>IκBASI</i>	1	16	12	15	52 (55%)	27	6 (22%)	21	12%
<i>CBP/PAG</i>	6	10	5	32	43 (48%)	16	7 (44%)	9	21%
<i>DOK3/DOK-L</i>	6	12	9	55	14 (16%)	11	5 (45%)	6	7%
<i>TAB2</i>	16	8	16	27	29 (36%)	10	2 (20%)	8	7%

a. A high PCR failure rate (internal control negative) was seen with *FANCG/XRCC9* prior to the introduction of improved protocols.

b. Percentage calculation denominator excludes PCR failures.

c. The percentage with no vector signal multiplied by the percentage with two FISH spots.

Fig. 14

REPLACEMENT SHEET

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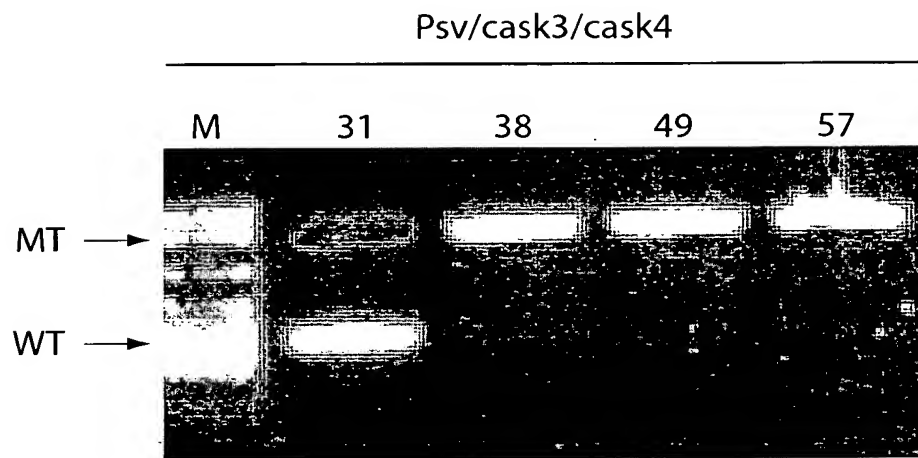


Fig. 15

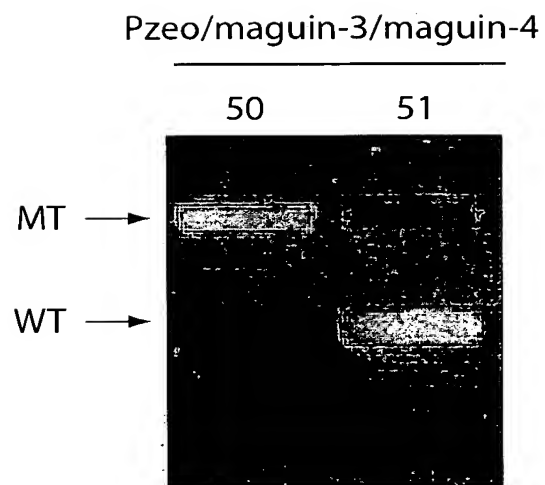


Fig. 16

REPLACEMENT SHEET

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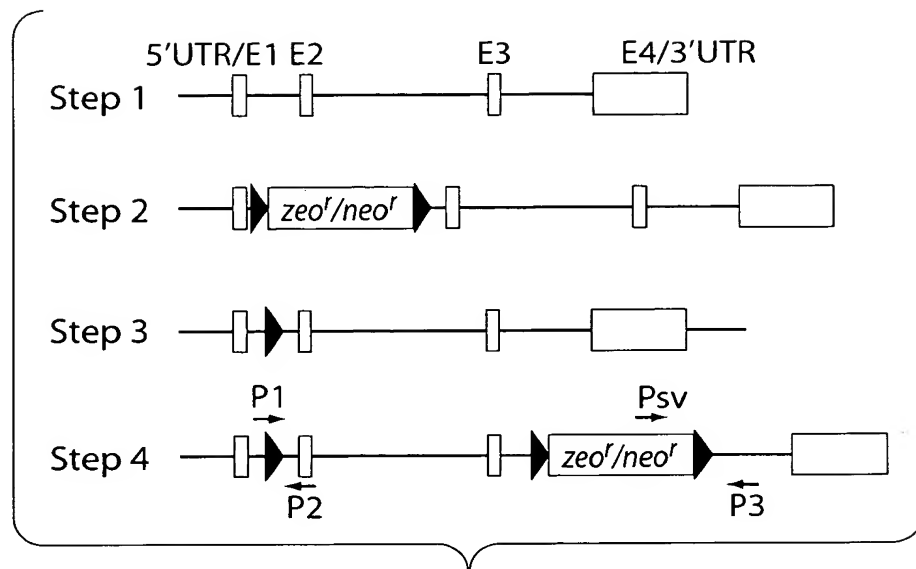


Fig. 17A

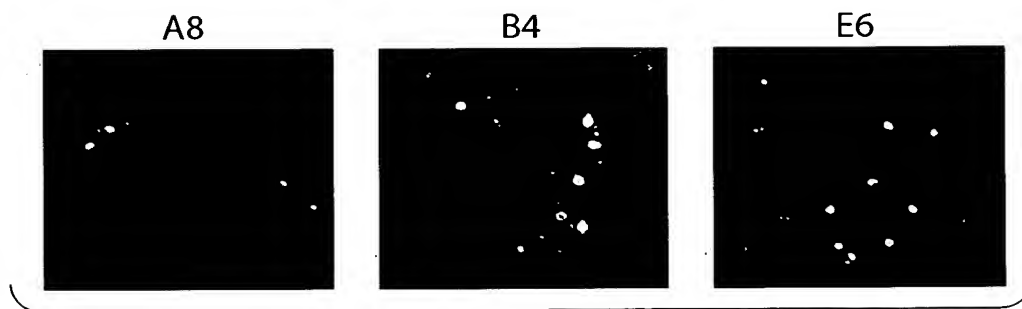


Fig. 17B

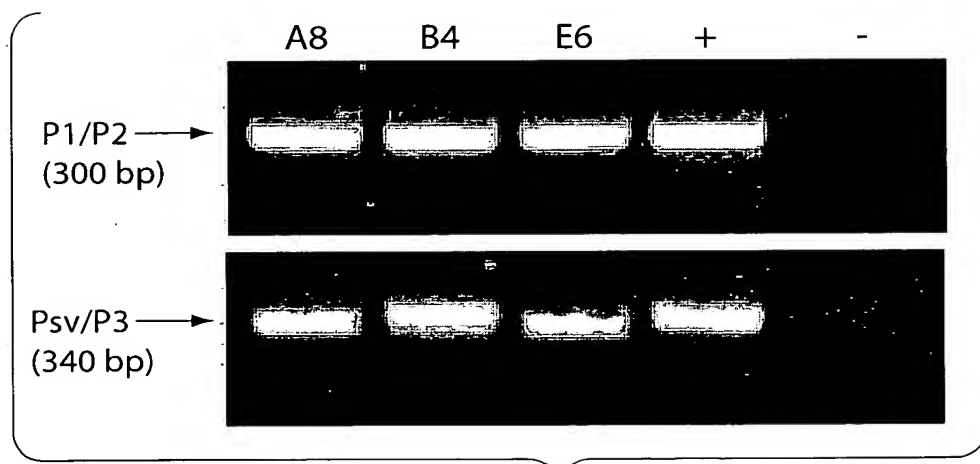


Fig. 17C